

IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/825,177

DATE: 09/01/2004 TIME: 13:05:26

Input Set : N:\Crf3\RULE60\10825177.raw Output Set: N:\CRF4\09012004\J825177.raw

```
1 <110> APPLICANT: ULLRICH, Axel
        NAYLER, Oliver
3 <120> TITLE OF INVENTION: CLK PROTEIN KINASES AND RELATED PRODUCTS AND METHODS
 4 <130> FILE REFERENCE: 038602/0431
 5 <140> CURRENT APPLICATION NUMBER: US/10/825,177
 6 <141> CURRENT FILING DATE: 2004-04-16
 7 <150> PRIOR APPLICATION NUMBER: US/09/905,999
8 <151> PRIOR FILING DATE: 2001-07-17
9 <150> PRIOR APPLICATION NUMBER: 09/127,248
10 <151> PRIOR FILING DATE: 1999-07-31
11 <150> PRIOR APPLICATION NUMBER: PCT/IB97/00946
12 <151> PRIOR FILING DATE: 1997-06-17
13 <150> PRIOR APPLICATION NUMBER: US 08/877,150
14 <151> PRIOR FILING DATE: 1997-06-17
15 <150> PRIOR APPLICATION NUMBER: US 60/034,286
16 <151> PRIOR FILING DATE: 1996-12-19
17 <160> NUMBER OF SEQ ID NOS: 26
18 <170> SOFTWARE: PatentIn version 3.0
20 <210> SEQ ID NO: 1
21 <211> LENGTH: 6
22 <212> TYPE: PRT
23 <213> ORGANISM: Mus musculus
24 <400> SEQUENCE: 1
25
        Asp Leu Lys Pro Glu Asn
26
        1
28 <210> SEQ ID NO: 2
29 <211> LENGTH: 6
30 <212> TYPE: PRT
31 <213> ORGANISM: Mus musculus
32 <400> SEQUENCE: 2
        Ala Met Met Glu Arg Ile
        1
36 <210> SEQ ID NO: 3
37 <211> LENGTH: 28
38 <212> TYPE: DNA
39 <213> ORGANISM: Mus musculus
40 <400> SEQUENCE: 3
         cgggatccct tcgccttgca gctttgtc
43 <210> SEQ ID NO: 4
44 <211> LENGTH: 30
45 <212> TYPE: DNA
```



28

47 <400> SEQUENCE: 4

46 <213> ORGANISM: Mus musculus

RAW SEQUENCE LISTING

DATE: 09/01/2004 PATENT APPLICATION: US/10/825,177 TIME: 13:05:26

48	cggaattcct agactgatac agtctgtaag	30
	SEQ ID NO: 5 LENGTH: 30	
	TYPE: DNA	
	ORGANISM: Mus musculus	
	SEQUENCE: 5	
55	tatageggee getagaetga tacagtetgt	30
	SEQ ID NO: 6	30
	LENGTH: 32	
	TYPE: DNA	
	ORGANISM: Mus musculus	
	SEQUENCE: 6	
62	_	32
	SEQ ID NO: 7	
	LENGTH: 39	
	TYPE: DNA	
	ORGANISM: Mus musculus	
	SEQUENCE: 7	
69	tatageggee geteacegae tgatateeeg aetggagte	39
	SEQ ID NO: 8	
	LENGTH: 30	
	TYPE: DNA	
	ORGANISM: Mus musculus	
	SEQUENCE: 8	
76	tccccgggg agacgatgca tcactgtaag	30
78 <210>	SEQ ID NO: 9	
	LENGTH: 39	
80 <212>	TYPE: DNA	
81 <213>	ORGANISM: Mus musculus	
82 <400>	SEQUENCE: 9	•
83	tatageggee gegetggeet geacetgtea tetgetggg	39
85 <210>	SEQ ID NO: 10	
86 <211>	LENGTH: 30	
87 <212>	TYPE: DNA	
88 <213>	ORGANISM: Mus musculus	
89 <400>	SEQUENCE: 10	
90	cggaattcat gcggcattcc aaacgaactc	30
92 <210>	SEQ ID NO: 11	
93 <211>	LENGTH: 39	
94 <212>	TYPE: DNA	
95 <213>	ORGANISM: Mus musculus	
96 <400>	SEQUENCE: 11	
97	tatageggee geeetgaete eeacteattt eetttttaa	39
	SEQ ID NO: 12	
	> LENGTH: 36	
	> TYPE: DNA .	
	> ORGANISM: Mus musculus	
	> SEQUENCE: 12	~ ~
104	cggaattccg ccaccatggc ccctatacta ggttat	36

RAW SEQUENCE LISTING

DATE: 09/01/2004 PATENT APPLICATION: US/10/825,177 TIME: 13:05:26

```
106 <210> SEQ ID NO: 13
107 <211> LENGTH: 36
108 <212> TYPE: DNA
109 <213> ORGANISM: Mus musculus
110 <400> SEQUENCE: 13
        gccaagettg ccaccatgge ecetataeta ggttat
                                                                                 36
113 <210> SEQ ID NO: 14
114 <211> LENGTH: 21
115 <212> TYPE: DNA
116 <213> ORGANISM: Mus musculus
117 <400> SEQUENCE: 14
118 gtagcagtaa gaatagttaa a
                                                                                 21
120 <210> SEQ ID NO: 15
121 <211> LENGTH: 24
122 <212> TYPE: DNA
123 <213> ORGANISM: Mus musculus
124 <400> SEQUENCE: 15
         gttgccctga ggatcattaa gaat
                                                                                 24
127 <210> SEQ ID NO: 16
128 <211> LENGTH: 24
129 <212> TYPE: DNA
130 <213> ORGANISM: Mus musculus
131 <400> SEQUENCE: 16
       gttgccctga ggatcatccg gaat
                                                                                 24
134 <210> SEQ ID NO: 17
135 <211> LENGTH: 30
136 <212> TYPE: DNA
137 <213> ORGANISM: Mus musculus
138 <400> SEQUENCE: 17
139 tacaattete aetgetaeat gtaageeate
                                                                                 30
141 <210> SEQ ID NO: 18
142 <211> LENGTH: 7
143 <212> TYPE: PRT
144 <213> ORGANISM: Artificial Sequence
145 <220> FEATURE:
146 <221> NAME/KEY: misc feature
147 <222> LOCATION: ()..()
148 <223> OTHER INFORMATION: Synthesized protein kinase
149 <400> SEQUENCE: 18
         His Arg Asp Leu Ala Ala Arg
151
153 <210> SEQ ID NO: 19
154 <211> LENGTH: 6
155 <212> TYPE: PRT
156 <213> ORGANISM: Artificial Sequence
157 <220> FEATURE:
158 <221> NAME/KEY: misc feature
159 <222> LOCATION: ()..()
160 <223> OTHER INFORMATION: Synthesized protein kinase
```

RAW SEQUENCE LISTING DATE: 09/01/2004
PATENT APPLICATION: US/10/825,177 TIME: 13:05:26

```
161 <221> NAME/KEY: misc feature
    162 <222> LOCATION: (2)..(2)
    163 <223> OTHER INFORMATION: Xaa at position 2 can be Val or Met
  > 164 <221> misc feature
    165 <222> LOCATION: (5)..(5)
    166 <223> OTHER INFORMATION: Xaa at position 5 can be Tyr or Phe
N--> 167 <400> 19
              Asp Xaa Trp Ser Xaa Gly
V--> 168
    169
    171 <210> SEQ ID NO: 20
    172 <211> LENGTH: 483
    173 <212> TYPE: PRT
    174 <213> ORGANISM: Mus musculus
    175 <400> SEQUENCE: 20
              Met Arg His Ser Lys Arg Thr Tyr Cys Pro Asp Trp Asp Glu Arg Asp
    176
    177
              Trp Asp Tyr Gly Thr Trp Arg Ser Ser Ser His Lys Arg Lys Lys
    178
    179
                                               25
    180
              Arg Ser His Ser Ser Ala Arg Glu Gln Lys Arg Cys Arg Tyr Asp His
              Ser Lys Thr Thr Asp Ser Tyr Tyr Leu Glu Ser Arg Ser Ile Asn Glu
    183
              Lys Ala Tyr His Ser Arg Arg Tyr Val Asp Glu Tyr Arg Asn Asp Tyr
    184
    185
                                   70
              Met Gly Tyr Glu Pro Gly His Pro Tyr Gly Glu Pro Gly Ser Arg Tyr
    186
    187
                               85
              Gln Met His Ser Ser Lys Ser Ser Gly Arg Ser Gly Arg Ser Ser Tyr
    188
    189
                           100
                                               105
              Lys Ser Lys His Arg Ser Arg His His Thr Ser Gln His His Ser His
    190
    191
              Gly Lys Ser His Arg Arg Lys Arg Ser Arg Ser Val Glu Asp Asp Glu
    192
    193
              Glu Gly His Leu Ile Cys Gln Ser Gly Asp Val Leu Ser Ala Arg Tyr
    194
    195
              Glu Ile Val Asp Thr Leu Gly Glu Gly Ala Phe Gly Lys Val Val Glu
    196
                                                   170
    197
              Cys Ile Asp His Lys Val Gly Gly Arg Arg Val Ala Val Lys Ile Val
    198
    199
                                               185
              Lys Asn Val Asp Arg Tyr Cys Glu Ala Ala Gln Ser Glu Ile Gln Val
    200
                                           200
    201
              Leu Glu His Leu Asn Thr Thr Asp Pro His Ser Thr Phe Arg Cys Val
    202
                                                            220
    203
                                    . 215
              Gln Met Leu Glu Trp Phe Glu His Arg Gly His Ile Cys Ile Val Phe
    204
                                                       235
    205
                                   230
              Glu Leu Leu Gly Leu Ser Thr Tyr Asp Phe Ile Lys Glu Asn Ser Phe
    206
                                                   250
    207
              Leu Pro Phe Arg Met Asp His Ile Arg Lys Met Ala Tyr Gln Ile Cys
    208
                                               265
    209
              Lys Ser Val Asn Phe Leu His Ser Met Lys Leu Thr His Thr Asp Leu
    210
```

RAW SEQUENCE LISTING DATE: 09/01/2004 PATENT APPLICATION: US/10/825,177 TIME: 13:05:26

211				275					280					285			
212		Lys	Pro	Glu	Asn	Ile	Leu	Phe	Val	Lys	Ser	Asp	Tyr	Thr	Glu	Ala	Tyr
213		-	290					295					300				
214		Asn		Lys	Met	Lvs	Ara	Asp	Glu	Ara	Thr	Ile	Val	Asn	Pro	Asp	Ile
215		305	110	270		272	310			5		315				1	320
			77-7	77.7	7	Dha		Cor	ת ד ת	The	TT		7 an	C1.,	uic	Uic	
216		гуѕ	Val	Val.	Asp		GIY	ser	Ата	1111	_	АБР	Asp	GIU	1113		DCI
217				_		325				_	330	_				335	
218		Thr	Leu	Val	Ser	Thr	Arg	His	Tyr	Arg	Ala	Pro	GLu	Val		Leu	Ala
219					340					345					350		
220		Leu	Gly	Trp	Ser	Gln	Pro	Cys	Asp	Val	Trp	Ser	Ile	Gly	Cys	Ile	Leu
221				355					360					365			
222		Ile	Glu	Tyr	Tvr	Leu	Glv	Phe	Thr	Val	Phe	Pro	Thr	His	Asp	Ser	Arq
223			370	4	4		-	375					380		_		_
224		Glu		Leu	Δla	Met	Met		Δra	Tle	T.e.11	Glv		Len	Pro	Lvs	His
			1113	пси	ALU	I-IC C		GIU	1119	110	cu	395		шец			400
225		385	~ 7	~ 3	-	em1	390	.	70	7	m		TT	TT-1	7	7	
226		Met.	11e	Gln	гуѕ		arg	гуs	Arg	Arg		Pne	HIS	HIS	Asp		ьeu
227						405					410		_			415	
228		Asp	Trp	Asp	Glu	His	Ser	Ser	Ala	Gly	Arg	Tyr	Val	Ser	Arg	Arg	Cys
229				-	420					425					430		
230		Lys	Pro	Leu	Lys	Glu	Phe	Met	Leu	Ser	Gln	Asp	Ala	Glu	His	Glu	Phe
231		_		435					440					445			
232		Leu	Phe	Asp	Leu	Val	Glv	Lvs	Ile	Leu	Glu	Tyr	Asp	Pro	Ala	Lys	Arq
233			450				1	455				•	460			-	
		Tlo		Leu	Laco	C1.11	777		Twe	uic	Pro	Dho		Туг	Pro	T.e.11	T.vc
234		TIE	1111	цеu	цув	Gru	AIA	пец	цуз	117.0	110	LIIC	LIIC	- y -	110	LC G	-, -
225		100					470					17 C					4 Q O
235		465		1			470					475					480
236		Lys	His				470					475					480
236 238	<210>	Lys SEQ	ID I	10: 2	21		470					475					480
236 238	<210> <211>	Lys SEQ	ID I	10: 2	21		470					475					480
236 238 239		Lys SEQ LEN	ID 1	NO: 2 499	21		470					475					480
236 238 239 240	<211>	Lys SEQ LENG TYPI	ID 1 GTH: E: PI	NO: 2 499 RT		ıscu.						475					480
236 238 239 240 241	<211><212><213>	Lys SEQ LENG TYPE ORGA	ID 1 GTH: E: PI ANISI	NO: 2 499 RT M: Mu	ıs mı	ıscul						475					480
236 238 239 240 241 242	<211> <212>	Lys SEQ LENG TYPE ORGA SEQU	ID 1 GTH: E: PI ANISI JENCI	NO: 2 499 RT M: Mu E: 23	ıs mu L		lus	Tvr	His	Ser	Ser		Arq	Gly	Ser	Arq	
236 238 239 240 241 242 243	<211><212><213>	Lys SEQ LENG TYPI ORGA SEQU Met	ID 1 GTH: E: PI ANISI JENCI	NO: 2 499 RT M: Mu E: 23	ıs mu L	Arg	lus	Tyr	His	Ser			Arg	Gly	Ser		480 Gly
236 238 239 240 241 242 243	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met	ID 1 GTH: E: PI ANISM JENCI Pro	NO: 2 499 RT M: Mu E: 2 His	ıs mu L Pro	Arg 5	lus Arg			•	10	Glu	•			15	Gly
236 238 239 240 241 242 243 244 245	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met	ID 1 GTH: E: PI ANISM JENCI Pro	NO: 2 499 RT M: Mu E: 23	ıs mu l Pro Glu	Arg 5	lus Arg			Arg	10	Glu	•		Arg	15	Gly
236 238 239 240 241 242 243 244 245 246	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser	ID 1 GTH: E: PI ANISI JENCI Pro Tyr	NO: 2 499 RT M: Mu E: 23 His	ıs mu l Pro Glu 20	Arg 5 His	lus Arg Tyr	Gln	Ser	Arg 25	10 Lys	Glu His	Lys	Arg	Arg 30	15 Arg	Gly Ser
236 238 239 240 241 242 243 244 245 246 247	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser	ID 1 GTH: E: PI ANISI JENCI Pro Tyr	NO: 2 499 RT M: Mu E: 2: His His	ıs mu l Pro Glu 20	Arg 5 His	lus Arg Tyr	Gln	Ser Asp	Arg 25	10 Lys	Glu His	Lys	Arg Arg	Arg 30	15 Arg	Gly Ser
236 238 239 240 241 242 243 244 245 246	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg	ID 1 GTH: E: PI ANISM JENCM Pro Tyr Ser	NO: 2 499 RT M: Mu E: 2: His His Trp 35	ıs mu l Pro Glu 20 Ser	Arg 5 His Ser	lus Arg Tyr Ser	Gln Ser	Ser Asp 40	Arg 25 Arg	10 Lys Thr	Glu His Arg	Lys Arg	Arg Arg 45	Arg 30 Arg	15 Arg Arg	Gly Ser Glu
236 238 239 240 241 242 243 244 245 246 247	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg	ID 1 GTH: E: PI ANISM JENCM Pro Tyr Ser	NO: 2 499 RT M: Mu E: 2: His His	ıs mu l Pro Glu 20 Ser	Arg 5 His Ser	lus Arg Tyr Ser	Gln Ser	Ser Asp 40	Arg 25 Arg	10 Lys Thr	Glu His Arg	Lys Arg	Arg Arg 45	Arg 30 Arg	15 Arg Arg	Gly Ser Glu
236 238 239 240 241 242 243 244 245 246 247 248	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser Arg	ID I STH: E: PH ANISM JENCH Pro Tyr Ser Ser 50	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	ıs mu l Pro Glu 20 Ser	Arg 5 His Ser Val	lus Arg Tyr Ser Arg	Gln Ser Ser 55	Ser Asp 40 Arg	Arg 25 Arg Ser	10 Lys Thr Ser	Glu His Arg Tyr	Lys Arg Asp 60	Arg Arg 45 Asp	Arg 30 Arg His	15 Arg Arg Ser	Gly Ser Glu Ser
236 238 239 240 241 242 243 244 245 246 247 248 249 250	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser Arg	ID I STH: E: PH ANISM JENCH Pro Tyr Ser Ser 50	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	ıs mu l Pro Glu 20 Ser	Arg 5 His Ser Val	lus Arg Tyr Ser Arg	Gln Ser Ser 55	Ser Asp 40 Arg	Arg 25 Arg Ser	10 Lys Thr Ser	Glu His Arg Tyr	Lys Arg Asp 60	Arg Arg 45 Asp	Arg 30 Arg His	15 Arg Arg Ser	Gly Ser Glu Ser
236 238 239 240 241 242 243 244 245 246 247 248 249 250 251	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser Arg	ID I STH: E: PH ANISM JENCH Pro Tyr Ser Ser 50	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	ıs mu l Pro Glu 20 Ser	Arg 5 His Ser Val	lus Arg Tyr Ser Arg	Gln Ser Ser 55	Ser Asp 40 Arg	Arg 25 Arg Ser	10 Lys Thr Ser	Glu His Arg Tyr	Lys Arg Asp 60	Arg Arg 45 Asp	Arg 30 Arg His	15 Arg Arg Ser	Gly Ser Glu
236 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser Arg Asp	ID I GTH: E: PH ANISM JENCH Pro Tyr Ser Ser 50 Arg	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	ls mul Pro Glu 20 Ser His Leu	Arg 5 His Ser Val	lus Arg Tyr Ser Arg Asp 70	Gln Ser Ser 55 Arg	Ser Asp 40 Arg	Arg 25 Arg Ser Tyr	10 Lys Thr Ser Cys	Glu His Arg Tyr Gly 75	Lys Arg Asp 60 Ser	Arg Arg 45 Asp	Arg 30 Arg His	15 Arg Arg Ser Arg	Gly Ser Glu Ser Asn 80
236 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253	<211><212><213><400>	Lys SEQ LENG TYPI ORGA SEQU Met 1 Ser Arg Asp	ID I GTH: E: PH ANISM JENCH Pro Tyr Ser Ser 50 Arg	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	ls mul Pro Glu 20 Ser His Leu	Arg 5 His Ser Val Tyr	lus Arg Tyr Ser Arg Asp 70	Gln Ser Ser 55 Arg	Ser Asp 40 Arg	Arg 25 Arg Ser Tyr	10 Lys Thr Ser Cys	Glu His Arg Tyr Gly 75	Lys Arg Asp 60 Ser	Arg Arg 45 Asp	Arg 30 Arg His	15 Arg Arg Ser Arg	Gly Ser Glu Ser Asn 80
236 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp	ID I ETH: E: PH ANISM PRO Tyr Ser Ser 50 Arg	NO: 2 499 RT M: MM E: 2: His His Trp 35 Tyr Arg	IS MU Pro Glu 20 Ser His Leu	Arg 5 His Ser Val Tyr Asp 85	lus Arg Tyr Ser Arg Asp 70 Arg	Gln Ser Ser 55 Arg	Ser Asp 40 Arg Arg	Arg 25 Arg Ser Tyr	10 Lys Thr Ser Cys Tyr 90	Glu His Arg Tyr Gly 75 Tyr	Lys Arg Asp 60 Ser Asp	Arg Arg 45 Asp Tyr	Arg 30 Arg His Arg	15 Arg Arg Ser Arg Phe 95	Gly Ser Glu Ser Asn 80 Arg
236 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 254 255	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp	ID I ETH: E: PH ANISM PRO Tyr Ser Ser 50 Arg	NO: 2 499 RT M: Mu E: 2: His His Trp 35 Tyr	IS MU Pro Glu 20 Ser His Leu Arg	Arg 5 His Ser Val Tyr Asp 85	lus Arg Tyr Ser Arg Asp 70 Arg	Gln Ser Ser 55 Arg	Ser Asp 40 Arg Arg	Arg 25 Arg Ser Tyr Ala Asn	10 Lys Thr Ser Cys Tyr 90	Glu His Arg Tyr Gly 75 Tyr	Lys Arg Asp 60 Ser Asp	Arg Arg 45 Asp Tyr	Arg 30 Arg His Arg Ser	15 Arg Arg Ser Arg Phe 95	Gly Ser Glu Ser Asn 80 Arg
236 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 254 255 256	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp	ID I STH: E: PH ANISM PRO Tyr Ser Ser 50 Arg Tyr Ser	NO: 2 499 RT M: MM E: 2: His Trp 35 Tyr Arg Ser Tyr	IS MU Pro Glu 20 Ser His Leu Arg Glu 100	Arg 5 His Ser Val Tyr Asp 85 Tyr	lus Arg Tyr Ser Arg Asp 70 Arg	Gln Ser Ser 55 Arg Gly Arg	Ser Asp 40 Arg Arg Glu Glu	Arg 25 Arg Ser Tyr Ala Asn 105	10 Lys Thr Ser Cys Tyr 90 Ser	Glu His Arg Tyr Gly 75 Tyr Ser	Lys Arg Asp 60 Ser Asp	Arg 45 Asp Tyr Thr	Arg 30 Arg His Arg Ser 110	Arg Arg Ser Arg Phe 95 Gln	Gly Ser Glu Ser Asn 80 Arg
236 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 254 255 256 257	<211><212><213><400>	Lys SEQ LENG TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp	ID I STH: E: PH ANISM PRO Tyr Ser Ser 50 Arg Tyr Ser	NO: 2 499 RT M: MM E: 22 His His Trp 35 Tyr Arg Ser Tyr	IS MU Pro Glu 20 Ser His Leu Arg Glu 100	Arg 5 His Ser Val Tyr Asp 85 Tyr	lus Arg Tyr Ser Arg Asp 70 Arg	Gln Ser Ser 55 Arg Gly Arg	Ser Asp 40 Arg Arg Glu Glu Arg	Arg 25 Arg Ser Tyr Ala Asn 105	10 Lys Thr Ser Cys Tyr 90 Ser	Glu His Arg Tyr Gly 75 Tyr Ser	Lys Arg Asp 60 Ser Asp	Arg 45 Asp Tyr Thr Arg Ser	Arg 30 Arg His Arg Ser 110	Arg Arg Ser Arg Phe 95 Gln	Gly Ser Glu Ser Asn 80 Arg
236 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 254 255 256 257 258	<211><212><213><400>	Lys SEQ LENC TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp Gln Ser	ID I GTH: E: PI ANISM PRO Tyr Ser Ser Arg Tyr Ser Ser	NO: 2 499 RT M: MM E: 2: His His Trp 35 Tyr Arg Ser Tyr	IS MU Pro Glu 20 Ser His Leu Arg Glu 100 Arg	Arg 5 His Ser Val Tyr Asp 85 Tyr	lus Arg Tyr Ser Arg Asp 70 Arg His	Gln Ser Ser 55 Arg Gly Arg	Ser Asp 40 Arg Arg Glu Glu Arg 120	Arg 25 Arg Ser Tyr Ala Asn 105 Arg	10 Lys Thr Ser Cys Tyr 90 Ser Arg	Glu His Arg Tyr Gly 75 Tyr Ser Arg	Lys Arg Asp 60 Ser Asp Tyr	Arg 45 Asp Tyr Thr Arg Ser 125	Arg 30 Arg His Arg Asp Ser 110 Arg	Arg Arg Ser Arg Phe 95 Gln Thr	Gly Ser Glu Ser Asn 80 Arg Arg
236 238 239 240 241 242 243 244 245 246 247 248 250 251 252 253 254 255 256 257	<211><212><213><400>	Lys SEQ LENC TYPH ORGA SEQU Met 1 Ser Arg Asp 65 Asp Gln Ser	ID I GTH: E: PI ANISM PRO Tyr Ser Ser Arg Tyr Ser Ser	NO: 2 499 RT M: MM E: 22 His His Trp 35 Tyr Arg Ser Tyr	IS MU Pro Glu 20 Ser His Leu Arg Glu 100 Arg	Arg 5 His Ser Val Tyr Asp 85 Tyr	lus Arg Tyr Ser Arg Asp 70 Arg His	Gln Ser Ser 55 Arg Gly Arg	Ser Asp 40 Arg Arg Glu Glu Arg 120	Arg 25 Arg Ser Tyr Ala Asn 105 Arg	10 Lys Thr Ser Cys Tyr 90 Ser Arg	Glu His Arg Tyr Gly 75 Tyr Ser Arg	Lys Arg Asp 60 Ser Asp Tyr	Arg 45 Asp Tyr Thr Arg Ser 125	Arg 30 Arg His Arg Asp Ser 110 Arg	Arg Arg Ser Arg Phe 95 Gln Thr	Gly Ser Glu Ser Asn 80 Arg Arg

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 09/01/2004

PATENT APPLICATION: US/10/825,177

TIME: 13:05:27

Input Set : N:\Crf3\RULE60\10825177.raw Output Set: N:\CRF4\09012004\J825177.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:19; Xaa Pos. 2,5

VERIFICATION SUMMARY

DATE: 09/01/2004

PATENT APPLICATION: US/10/825,177

TIME: 13:05:27

Input Set : N:\Crf3\RULE60\10825177.raw Output Set: N:\CRF4\09012004\J825177.raw

:164 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19 :167 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19

:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0